

# MATH 598: TOPICS IN STATISTICS

## BAYESIAN NONPARAMETRIC METHODS

The *Dirichlet Process* is a probability distribution on the space of discrete distributions on  $\mathbb{R}$  characterized by

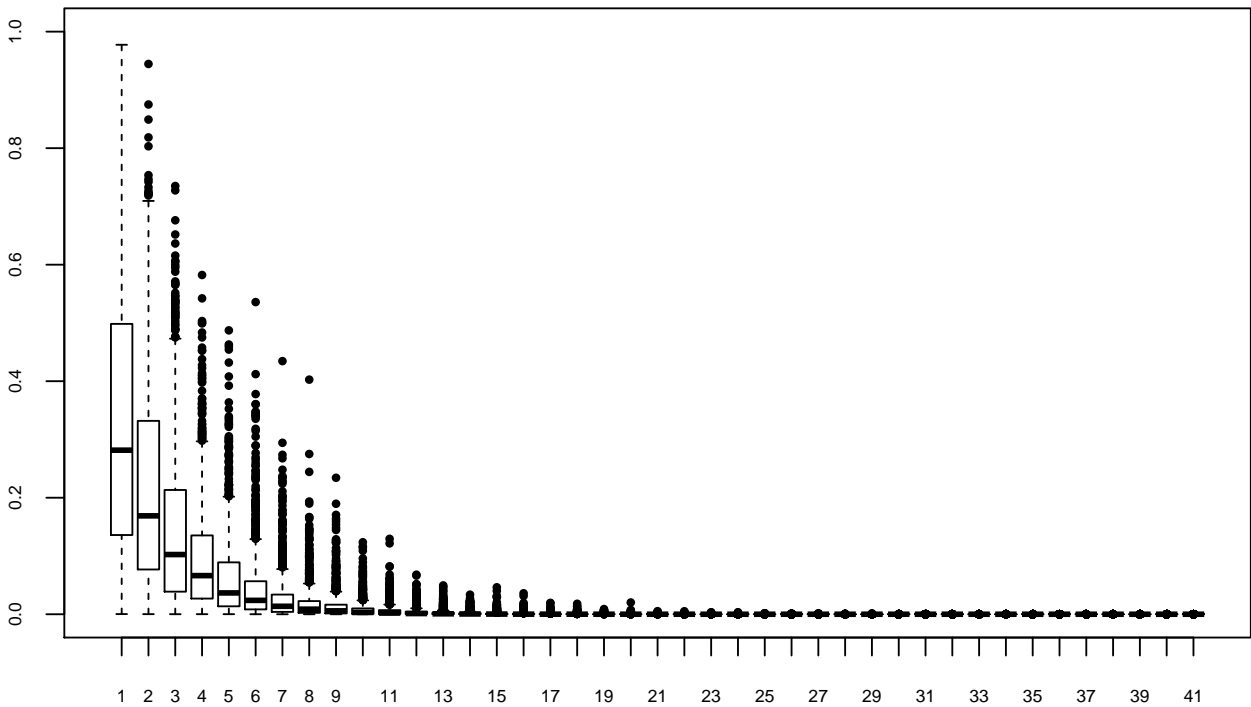
- a base measure,  $G_X$ , which is some probability measure on  $\mathbb{R}$ , from which are drawn a countable collection of random variables  $X_1, X_2, \dots$
- a countable collection of random variables  $\pi_1, \pi_2, \dots$  with  $\pi_i > 0$  such that  $\sum_{i=1}^{\infty} \pi_i = 1$  that is constructed in the following way:  $\pi_1 = V_1 \sim \text{Beta}(1, \alpha)$  and

$$\pi_i = V_i \prod_{j=1}^{i-1} (1 - V_j)$$

where for each  $j$ ,  $V_j \sim \text{Beta}(1, \alpha)$  for some parameter  $\alpha > 0$ . Note that by construction, in expectation  $\pi_1 > \pi_2 > \dots$ , which facilitates truncation at some finite value  $M$ .

```
M<-41
alpha<-2
pi.mat<-matrix(0,nrow=1000,ncol=M)
for(i in 1:1000){
  V<-rbeta(M,1,alpha)
  pi.mat[i,]<-c(V[1],cumprod(1-V[-M])*V[-1])
}
par(mar=c(2,2,3,0))
boxplot(pi.mat,pch=19,cex=0.5,ylim=range(0,1),cex.axis=0.6)
title(expression(paste('Boxplot of stick-breaking draws of ', pi)),line=1)
```

Boxplot of stick-breaking draws of  $\pi$

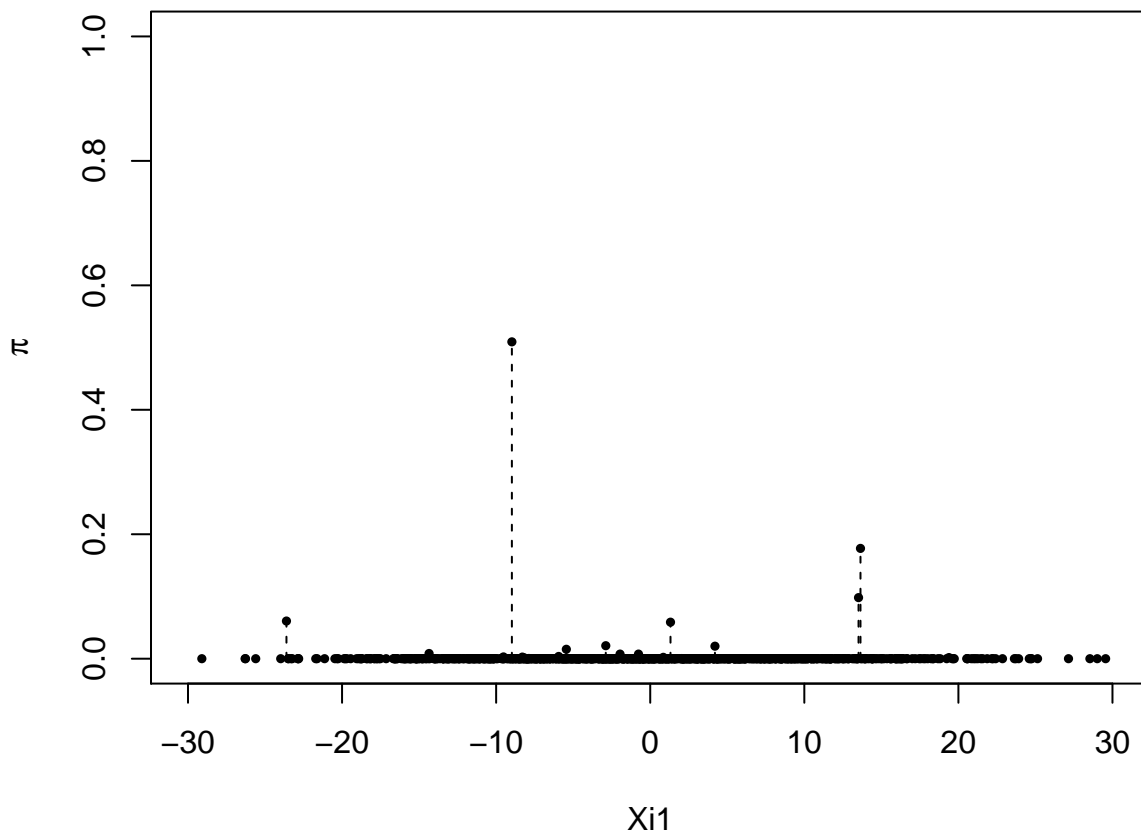


The collections  $\{X_1, X_2, \dots\}$  and  $\{\pi_1, \pi_2, \dots\}$  define a random discrete distribution with mass function  $\tilde{f}$  where

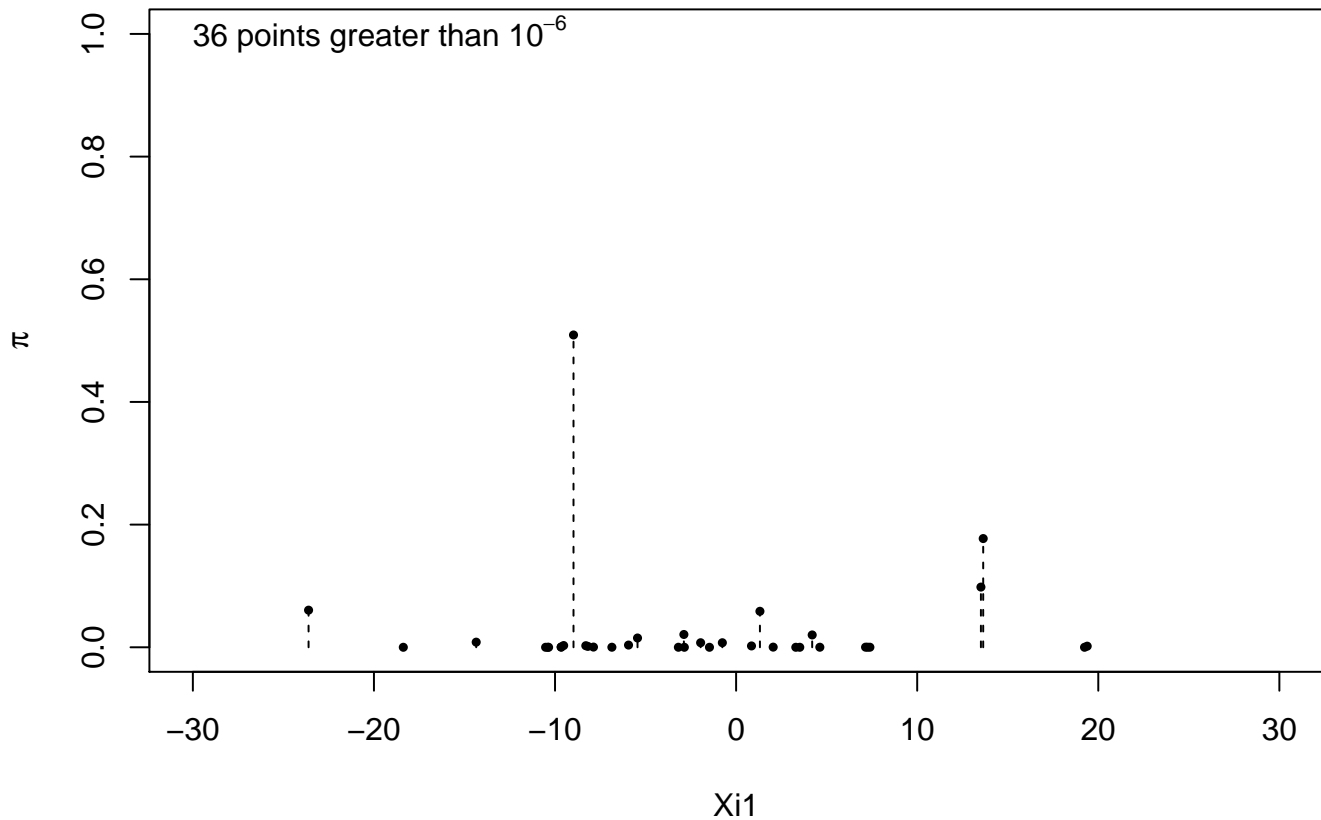
$$\tilde{f}(x) = \sum_{i=1}^{\infty} \pi_i \delta_{X_i}(x)$$

where  $\delta_{x_i}(x) = 1 \iff x = x_i$ . We write  $\tilde{f} \sim DP(\alpha, G_X)$  to denote that  $\tilde{f}$  is drawn from the Dirichlet Process.

```
M<-1001
alpha<-2
set.seed(1010)
#Stick-breaking
Xi<-rnorm(M,0,10) #Base measure
V<-rbeta(M,1,alpha)
pi.vec<-c(V[1],cumprod(1-V[-M])*V[-1])
Xi1<-Xi
pi1<-pi.vec
par(mar=c(4,4,1,0))
plot(Xi1,pi1,type="n",ylim=range(0,1),ylab=expression(pi),xlim=range(-30,30))
points(Xi1,pi1,pch=19,xlim=range(-30,30),cex=0.5)
for(i in 1:M){lines(c(Xi1[i],Xi1[i]),c(0,pi1[i]),lty=2)}
```



```
par(mar=c(4,4,1,0))
plot(Xi1,pi1,type="n",ylim=range(0,1),ylab=expression(pi),xlim=range(-30,30))
points(Xi[pi1>1e-6],pi.vec[pi1>1e-6],pch=19,xlim=range(-30,30),cex=0.5)
for(i in 1:M){
  if(pi1[i] < 1e-6) next
  lines(c(Xi1[i],Xi1[i]),c(0,pi1[i]),lty=2)
}
nval<-sum(pi1>1e-6)
text(-30,1,substitute(paste(nv, ' points greater than ', 10^{-6}),list(nv=nval)),adj=0)
```



```

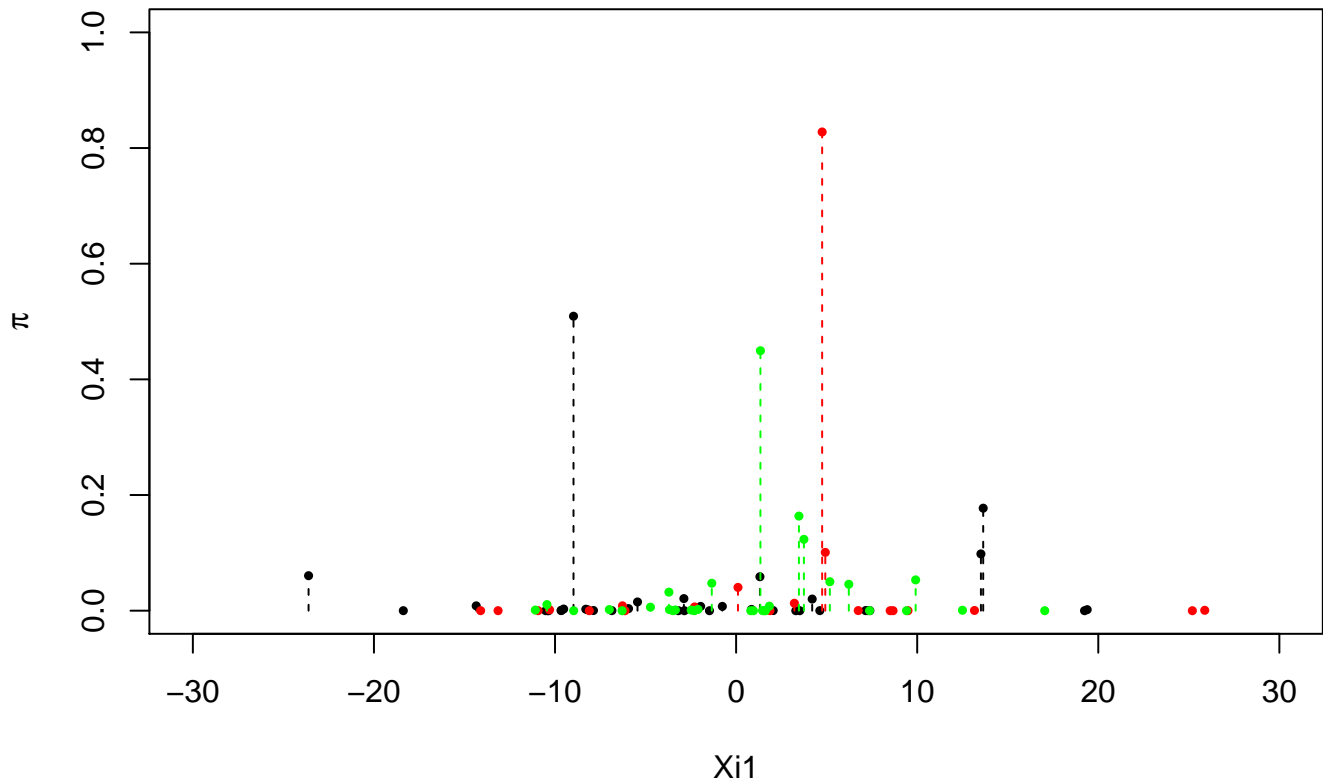
Xi<-rnorm(M,0,10)
V<-rbeta(M,1,alpha)
pi.vec<-c(V[1],cumprod(1-V[-M])*V[-1])
Xi2<-Xi
pi2<-pi.vec

Xi<-rnorm(M,0,10)
V<-rbeta(M,1,alpha)
W<-1-V
pi.vec<-c(V[1],cumprod(1-V[-M])*V[-1])
Xi3<-Xi
pi3<-pi.vec

par(mar=c(4,4,2,0))
plot(Xi1,pi1,type="n",ylim=range(0,1),ylab=expression(pi),xlim=range(-30,30))
title(expression(paste('Three draws from the ',DP(alpha,G[X])))
points(Xi1[pi1>1e-6],pi1[pi1>1e-6],pch=19,xlim=range(-30,30),cex=0.5)
for(i in 1:M){
  if(pi1[i] < 1e-6) next
  lines(c(Xi1[i],Xi1[i]),c(0,pi1[i]),lty=2)
}
points(Xi2[pi2>1e-6],pi2[pi2>1e-6],pch=19,xlim=range(-30,30),col="red",cex=0.5)
for(i in 1:M){
  if(pi2[i] < 1e-6) next
  lines(c(Xi2[i],Xi2[i]),c(0,pi2[i]),lty=2,col="red")
}
points(Xi3[pi3>1e-6],pi3[pi3>1e-6],pch=19,xlim=range(-30,30),col="green",cex=0.5)
for(i in 1:M){
  if(pi3[i] < 1e-6) next
  lines(c(Xi3[i],Xi3[i]),c(0,pi3[i]),lty=2,col="green")
}

```

### Three draws from the DP( $\alpha, G_X$ )



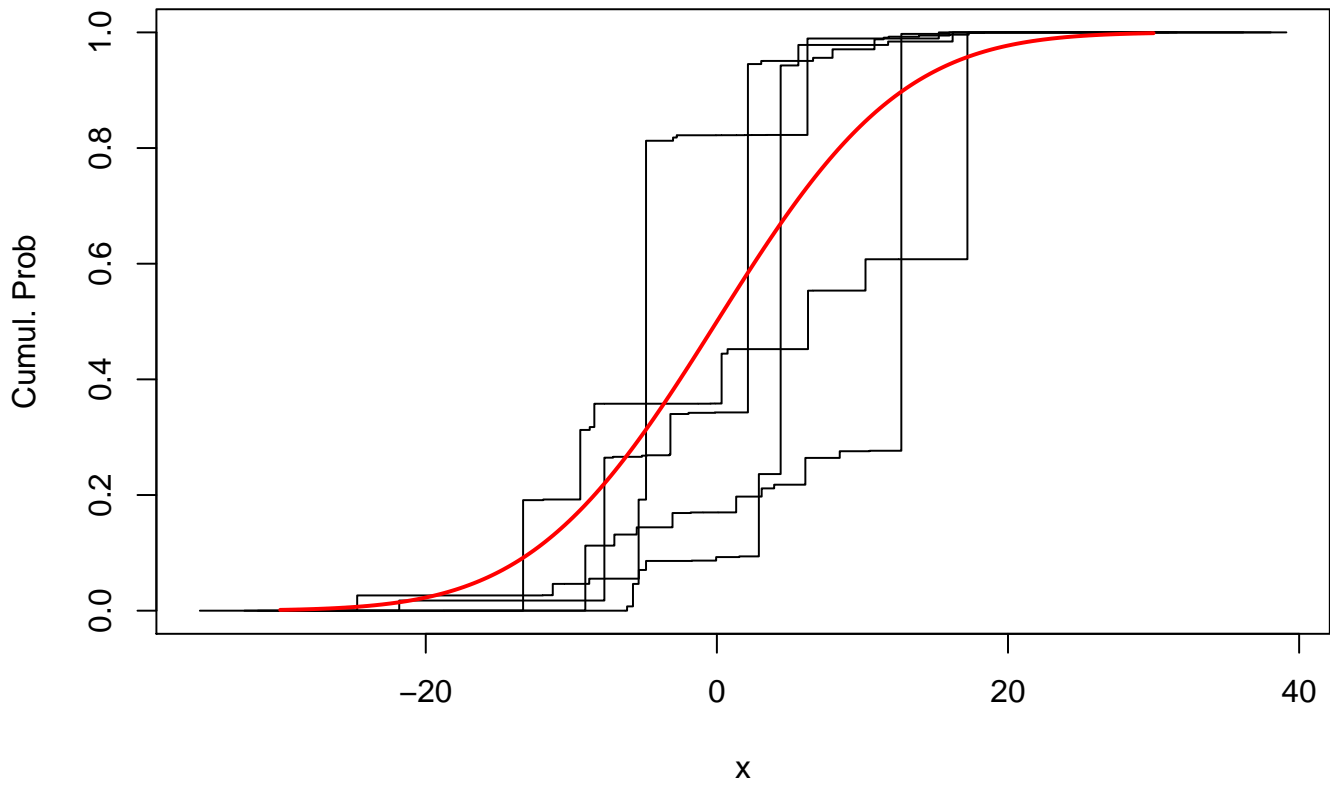
```

Xi<-rnorm(M,0,10)
V<-rbeta(M,1,alpha)
pi.vec<-c(V[1],cumprod(1-V[-M])*V[-1])
XiLarge<-Xi
piLarge<-pi.vec

xi.sort<-XiLarge[order(XiLarge)]
pi.sort<-pi.vec[order(XiLarge)]
Pi<-cumsum(pi.sort)
par(mar=c(4,4,2,0))
plot(xi.sort,Pi,type="s",ylim=range(0,1),xlab="x",ylab="Cumul. Prob")
title(expression(paste('Five replicate draws of the random cdf ',(alpha==2))))
for(i in 1:4){
  Xi<-rnorm(M,0,10)
  V<-rbeta(M,1,alpha)
  pi.vec<-c(V[1],cumprod(1-V[-M])*V[-1])
  XiLarge<-Xi
  piLarge<-pi.vec
  xi.sort<-XiLarge[order(XiLarge)]
  pi.sort<-pi.vec[order(XiLarge)]
  Pi<-cumsum(pi.sort)
  lines(xi.sort,Pi,type="s")
}
xv<-seq(-30,30,length=1001)
yv<-pnorm(xv,0,10)
lines(xv,yv,col="red",lwd=2)

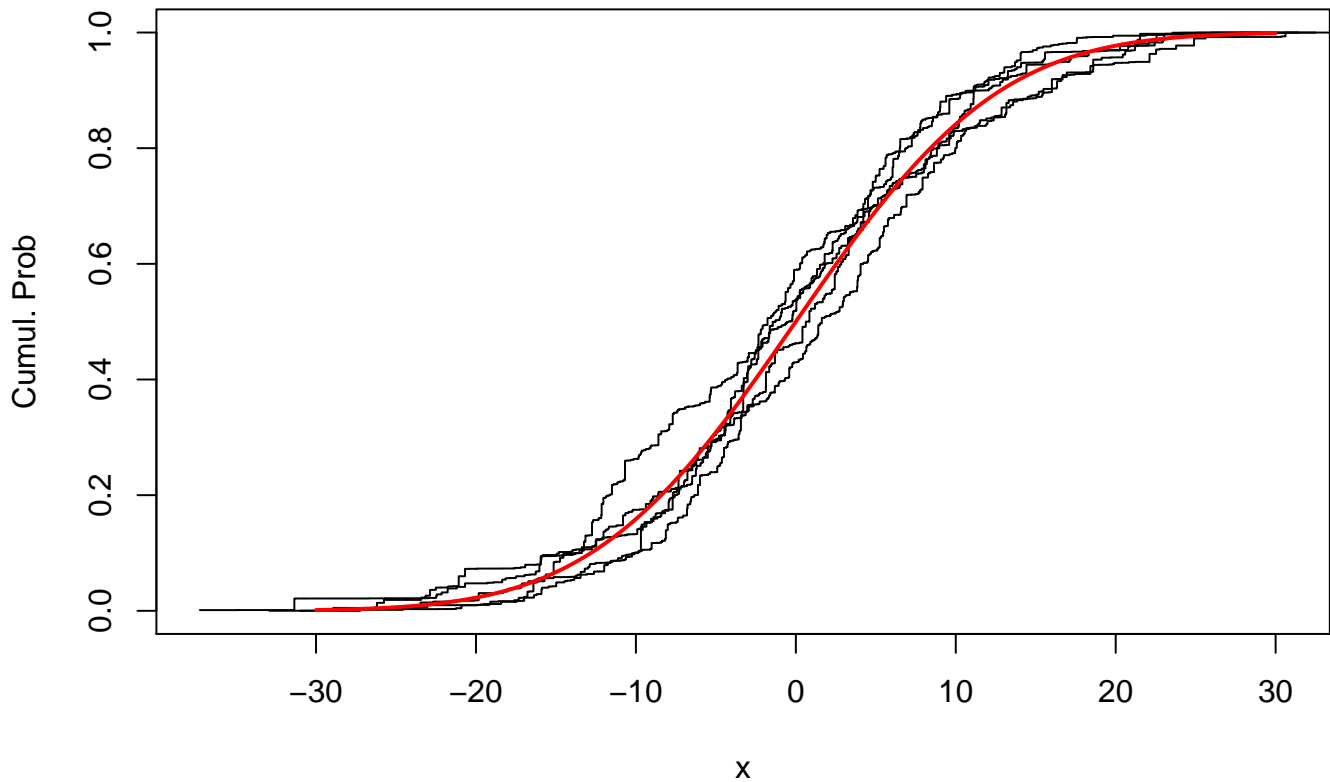
```

Five replicate draws of the random cdf ( $\alpha = 2$ )



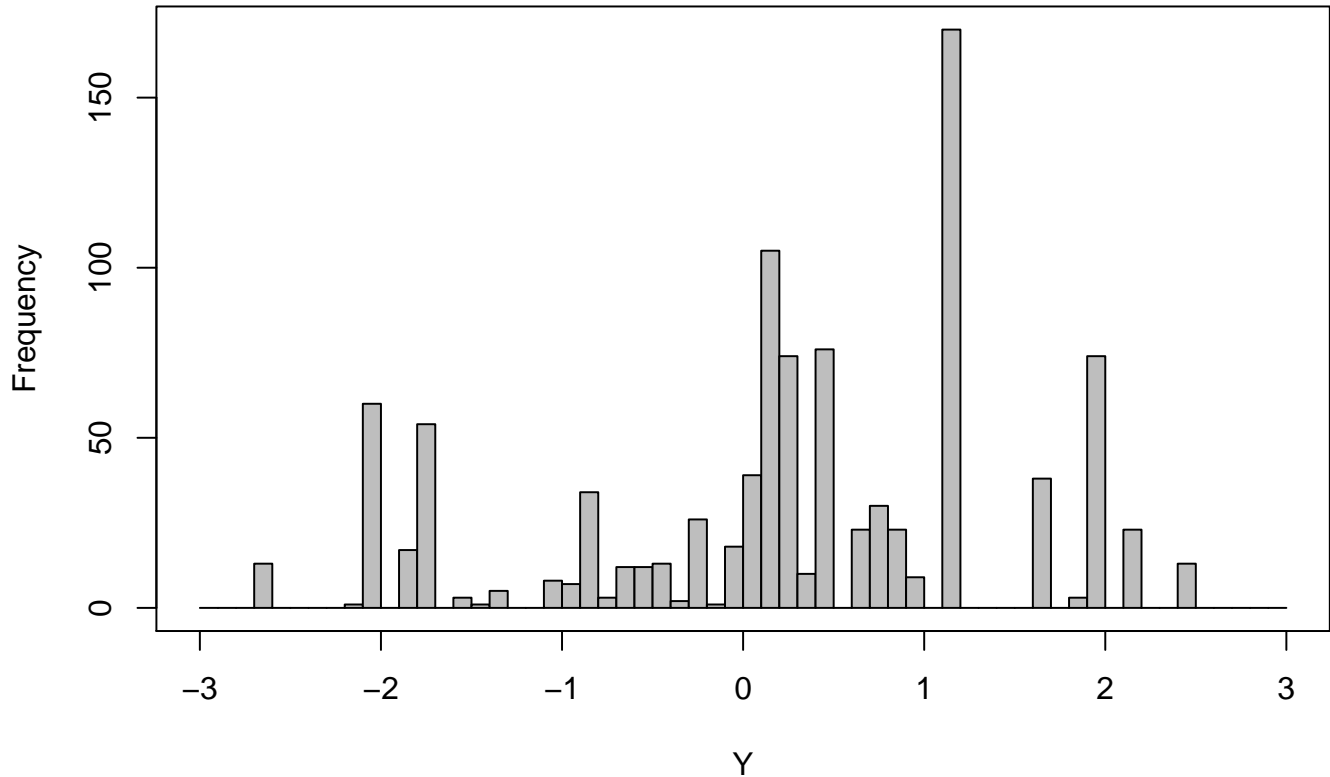
The parameter  $\alpha$  is a precision parameter; as  $\alpha$  increases, the random distribution becomes more concentrated on the base measure.

Five replicate draws of the random cdf ( $\alpha = 100$ )



The random distribution  $\tilde{f}$  can be sampled to produce iid data  $Y_1, \dots, Y_n$  in a straightforward fashion. In the simulation below, we simulate  $\tilde{f}$  with  $\alpha = 10$  and  $G_X \equiv Normal(0, 1)$

Sample of data from  $\tilde{f}$



Note that in the sample of size  $n = 1000$ , there are only 64 distinct  $Y$  values

```
tabY<-table(Y)
names(tabY)<-as.character(round(as.numeric(names(tabY)),5))
tabY
```

+	-2.63151	-2.16276	-2.08518	-1.81156	-1.75768	-1.70421	-1.54031	-1.40852
+	13	1	60	17	45	9	3	1
+	-1.32673	-1.09304	-0.90964	-0.86634	-0.83135	-0.78091	-0.73734	-0.69322
+	5	8	7	1	33	2	1	1
+	-0.66846	-0.62998	-0.53459	-0.50782	-0.47949	-0.41675	-0.33048	-0.32752
+	8	3	11	1	12	1	1	1
+	-0.29525	-0.28987	-0.26925	-0.1323	-0.09247	-0.08415	-0.0704	-0.05806
+	1	4	21	1	1	6	5	5
+	-0.0233	0.01556	0.10959	0.18201	0.21553	0.2375	0.25179	0.25219
+	1	39	104	1	4	13	1	8
+	0.25484	0.28758	0.31203	0.3336	0.3526	0.38624	0.41278	0.49196
+	1	47	5	1	3	1	69	7
+	0.60717	0.61285	0.65872	0.70102	0.71598	0.72779	0.81569	0.94265
+	11	9	3	2	2	26	23	9
+	1.11499	1.13418	1.60025	1.68618	1.85047	1.99737	2.13939	2.43685
+	132	38	3	35	3	74	23	13

and thus we observe *clustering*: this is due to the fact that  $\tilde{f}$  is discrete.

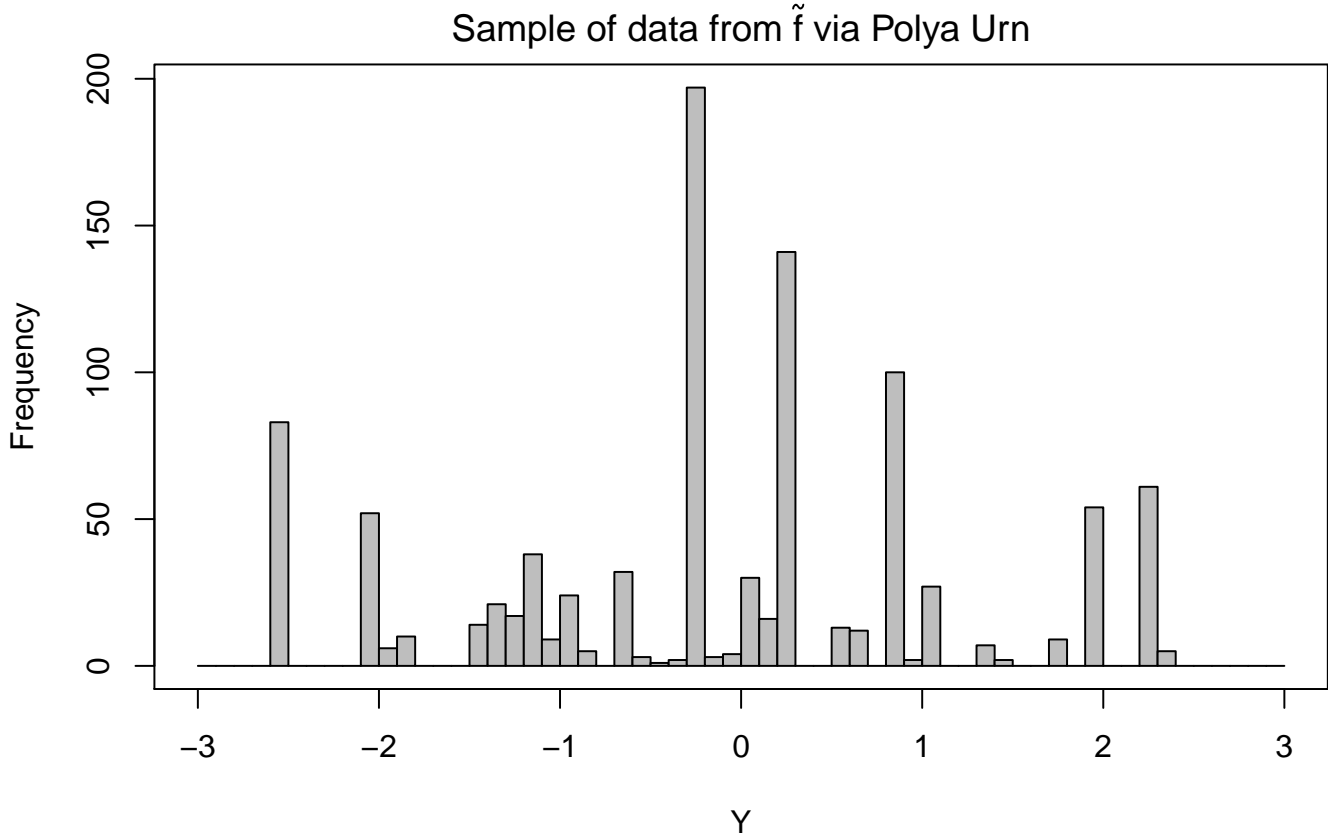
We can also use a *Polya Urn* scheme to sample these data: this recursive scheme proceeds as follows:

1. simulate  $Y_1 \sim G_X$ ;
2. for  $i = 2, 3, \dots, n$  generate  $Y_i$  from the mixture distribution

$$\frac{\alpha}{\alpha + i - 1} G_X(\cdot) + \frac{1}{\alpha + i - 1} \sum_{j=1}^{i-1} \delta_{Y_j}(\cdot)$$

that is

- with probability  $\alpha/(\alpha + i - 1)$ , simulate  $Y_i \sim G_X(\cdot)$ ;
- with probability  $1/(\alpha + i - 1)$ , simulate  $Y_i$  uniformly on  $\{Y_1, \dots, Y_{i-1}\}$ .



Here there are 49 distinct  $Y$  values. The Polya Urn method reconstructs a sample of exchangeable observables according to the de Finetti representation

$$f_{Y_1, \dots, Y_n}(y_1, \dots, y_n) = \int \prod_{i=1}^n \tilde{f}(y_i) \pi_0(d\tilde{f})$$

by sampling directly from the left hand side using the factorization

$$f_{Y_1, \dots, Y_n}(y_1, \dots, y_n) = f_{Y_1}(y_1) \prod_{i=2}^n f_{Y_i|Y_1, \dots, Y_{i-1}}(y_i|y_1, \dots, y_{i-1})$$

where

$$f_{Y_i|Y_1, \dots, Y_{i-1}}(y_i|y_1, \dots, y_{i-1}) = \frac{\alpha}{\alpha + i - 1} g_X(y_i) + \frac{1}{\alpha + i - 1} \sum_{j=1}^{i-1} \delta_{Y_j}(y_i)$$

where  $g_X(\cdot)$  is the density corresponding to  $G_X(\cdot)$ .

In a *Dirichlet Process Mixture* model we add another stage that brings in a continuous distribution. For example, could treat each  $x_i$  as the location of a normal density, and consider generating a  $y$  for each

$$\begin{aligned}
 x_1, x_2, \dots &\sim G_X \\
 \pi_1, \pi_2, \dots &\text{ generated by stick-breaking.} \\
 y_i &\sim \phi((y_i - x_i)/\sigma) \quad i = 1, 2, \dots
 \end{aligned}$$

Then the random density function generated takes the form

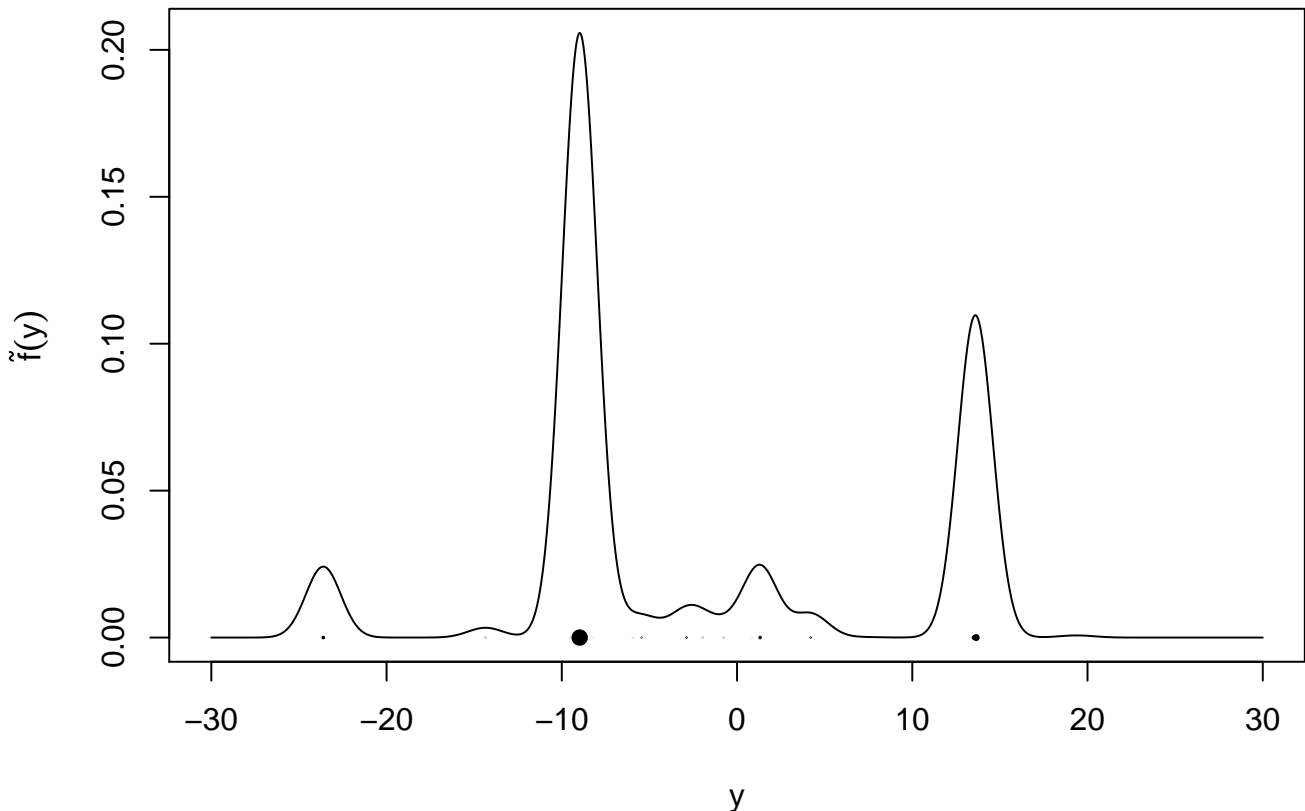
$$\tilde{f}(y) = \sum_{i=1}^{\infty} \pi_i \phi((y - x_i)/\sigma)$$

that is, an **infinite mixture model**. In the plot below we have  $\alpha = 2$ , with  $G_X(x) \equiv Normal(0, \lambda^2)$  with  $\lambda = 10$ .

```

set.seed(1010)
M<-1001
alpha<-2
sigma<-1;lambda<-10
Xi<-rnorm(M,0,lambda)
V<-rbeta(M,1,alpha)
pi.vec<-c(V[1],cumprod(1-V[-M]))*V[-1])
xv<-seq(-30,30,by=0.01)
yv<-xv*0
for(i in 1:M){
  yv<-yv+pi.vec[i]*dnorm(xv,Xi[i],sigma)
}
par(mar=c(4,5,2,0))
plot(xv,yv,type='l',xlab='y',ylab=expression(widetilde(f)(y)))
points(Xi,Xi*0,cex=2*pi.vec,pch=19)

```





**Gibbs sampler:** For fully Bayesian inference consider the de Finetti construction

- a prior model for  $f$  that is  $DPM(\alpha, G_X, g_Y; \theta)$  where  $\theta$  represents the parameters that appear in  $G_X$  and  $g_Y$ .
- conditional on  $f$ , data  $y_1, y_2, \dots, y_n \sim f$  independently

We wish to compute the posterior for  $f$ . We use the hierarchical formulation

$$\begin{aligned} y_j | x_j &\stackrel{\text{ind.}}{\sim} g_Y(y_j | x_j; \theta) & j = 1, \dots, n \\ x_1, \dots, x_n &\sim DP(\alpha, G_X; \theta) \\ \theta &\sim \pi_0(\theta). \end{aligned}$$

The latent variables  $x_1, \dots, x_n$  are also treated as parameters. They can be sampled using an MCMC **Gibbs sampler** scheme. For  $j = 1, \dots, n$ , we sample

$$x_j | \mathbf{x}_{(j)}, \mathbf{y} \sim w_0 p(x_j | y_j) + \sum_{l \neq j} w_l \delta_{x_l}$$

where

- $\mathbf{y} = (y_1, \dots, y_n)^\top$
- $\mathbf{x}_{(j)} = (x_1, \dots, x_{j-1}, x_{j+1}, \dots, x_n)^\top$ .
- $w_0$  is proportional to  $\alpha$  times the **prior predictive** of  $y_j$
- $w_l$  is proportional to the **likelihood** of  $y_j | x_l$
- $p(x_j | y_j)$  is the **posterior** for  $x_j$  given  $y_j$ .

**Marginalized sampler:** To be more efficient, we can use the **clustering** property: suppose that at a given iteration of the MCMC, there are  $K$  clusters labelled 1 to  $K$ , where  $K \leq n$ . Label the  $K$  distinct  $x$  values  $z_1, \dots, z_K$  and for each  $j$ , define the corresponding cluster label  $c_j$  where

$$c_j = k \quad \iff \quad x_j = z_k$$

We can update the  $c_j$ s instead of the  $x_j$ s which will be more computationally efficient; we are clustering  $x$ s to the **cluster centres** at the  $z$  values. For  $i = 1, \dots, n$ , let

- $n_1(i), \dots, n_K(i)$  denote the number of items in clusters 1,  $\dots$ ,  $K$
- $\mathbf{y}_1(i), \dots, \mathbf{y}_K(i)$  denote the vectors of  $y$  values currently allocated to the  $K$  clusters

**if the  $i$ th data point is removed.** For  $i = 1, \dots, n$ , we sample the cluster labels in a Gibbs sampler with conditional probabilities

$$\Pr[c_i = k | c_{(i)}] \propto \frac{n_k(i)}{\alpha + n - 1} p(y_i | \mathbf{y}_k(i)) \quad k = 1, \dots, K$$

and

$$\Pr[c_i = K + 1 | c_{(i)}] \propto \frac{\alpha}{\alpha + n - 1} p(y_i)$$

In this expression

- $p(y_i | \mathbf{y}_k(i))$  is the **posterior predictive** density for  $y_i$ , assuming that  $y_i$  comes from cluster  $k$ .
- $p(y_i)$  is the **prior predictive** density for  $y_i$ , assuming that  $y_i$  comes from a **new cluster** not currently represented in the data.

In this formulation, we have **integrated out** the Dirichlet Process analytically. Thus we can simply sample the cluster labels in turn, and then sample the  $z_1, \dots, z_K$  values; this will allow us to do density estimation. By the usual calculation

$$p(y_i | \mathbf{y}_k(i)) = \int g_Y(y_i | x) p(x | \mathbf{y}_k(i)) dx$$

where

$$p(x | \mathbf{y}_k(i)) \propto p(\mathbf{y}_k(i) | x) p(x) = \left\{ \prod_{l \neq i} g_Y(y_l | x) \right\} p(x)$$

gives the posterior distribution for the  $k$ th cluster centre. Similarly

$$p(y_i) = \int g_Y(y_i | x)p(x) dx$$

In the earlier Normal model, suppose for simplicity that  $G_X$  is the  $Normal(0, \lambda^2)$  density:

$$x_i \sim Normal(0, \lambda^2) \quad y_i | x_i \sim Normal(x_i, \sigma^2)$$

Then by standard calculations

$$p(y_i | \mathbf{y}_k(i)) \equiv Normal \left( \frac{n_k(i)\bar{y}_k(i)/\sigma^2}{n_k(i)/\sigma^2 + 1/\lambda^2}, \frac{(n_k(i) + 1)/\sigma^2 + 1/\lambda^2}{n_k(i)/\sigma^2 + 1/\lambda^2} \sigma^2 \right)$$

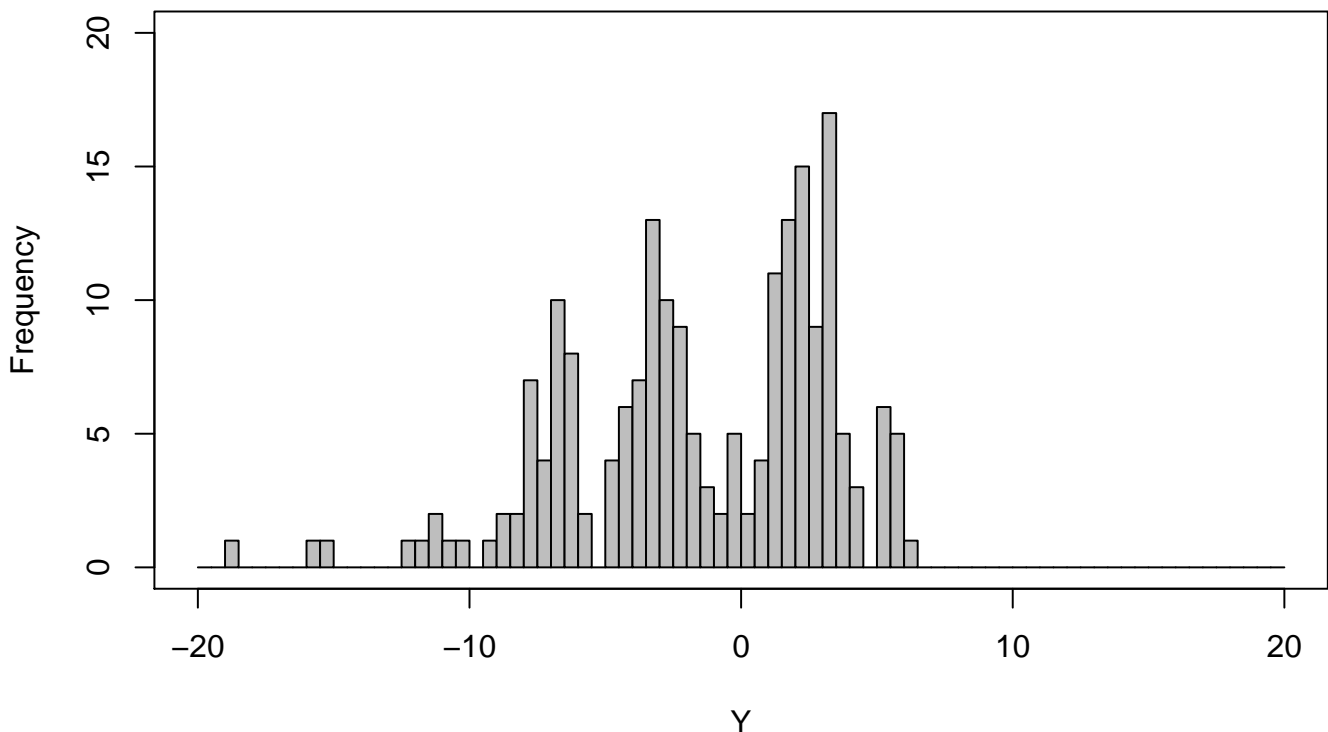
and

$$p(y_i) \equiv Normal(0, \sigma^2 + \lambda^2).$$

The marginalized version of the algorithm often mixes more quickly than the Gibbs sampler on the cluster centres.

```
set.seed(1091)
n<-200
alpha<-2
sigma<-1; lambda<-10
X<-numeric(n)
X[1]<-rnorm(1,0,lambda)
for(j in 2:n){
  u<-runif(1)
  if(u < alpha/(alpha+j-1)){
    X[j]<-rnorm(1,0,lambda)
  }else{
    X[j]<-sample(X[1:(j-1)],size=1)
  }
}
Y<-rnorm(n,X,sigma)
par(mar=c(4,4,2,0))
hist(Y,breaks=seq(-20,20,by=0.50),ylim=range(0,20),col='gray',main='Simulated data');box()
```

**Simulated data**



```

True.K<-length(table(X))

#####
set.seed(1010)
index<-c(1:n)
Kval<-5
Yq<-quantile(Y,prob=c(0:Kval)/Kval)
Cvec<-as.numeric(cut(Y,Yq,include.lowest=T))
Ysum.Vec<-rep(0,Kval)
for(k in 1:Kval){Ysum.Vec[k]<-sum(Y[Cvec==k])}
Nvec<-as.numeric(table(Cvec))

nreps<-10000
Nsamp<-rep(0,nreps)
yv<-seq(-20,20,by=0.01)
fyv.samp<-numeric()

for(irep in 1:nreps){
  for(i in 1:n){
    sum.Vec<-Ysum.Vec
    sum.Vec[Cvec[i]]<-sum.Vec[Cvec[i]]-Y[i]
    nvec<-Nvec
    nvec[Cvec[i]]<-nvec[Cvec[i]]-1
    if(nvec[Cvec[i]] > 0){
      mvec<-sum.Vec/nvec
      Mvec<-(sum.Vec/sigma^2)/(nvec/sigma^2+1/lambda^2)
      Svec<-sigma^2*((nvec+1)/sigma^2+1/lambda^2)/(nvec/sigma^2+1/lambda^2)
      pvec<-nvec*dnorm(rep(Y[i],Kval),Mvec,sqrt(Svec))
      pvec<-c(pvec,alpha*dnorm(Y[i],0,sqrt(sigma^2 + lambda^2)))/(n-1+alpha)
      pvec<-pvec/sum(pvec)
      Ksamp<-sample(c(1:(Kval+1)),size=1,prob=pvec)
      Cvec[i]<-Ksamp
      if(Ksamp == Kval+1){
        Kval<-Kval+1
        Ysum.Vec<-c(sum.Vec,Y[i])
        Nvec<-c(nvec,1)
      }else{
        Ysum.Vec<-sum.Vec
        Ysum.Vec[Ksamp]<-sum.Vec[Ksamp]+Y[i]
        Nvec<-nvec
        Nvec[Ksamp]<-nvec[Ksamp]+1
      }
    }else{
      Ktmp<-Cvec[i]
      sum.Vec<-sum.Vec[-Ktmp]
      nvec<-nvec[-Ktmp]
      Kval<-Kval-1
      Cvec[Cvec > Ktmp]<-Cvec[Cvec > Ktmp]-1

      mvec<-sum.Vec/nvec
      Mvec<-(sum.Vec/sigma^2)/(nvec/sigma^2+1/lambda^2)
      Svec<-sigma^2*((nvec+1)/sigma^2+1/lambda^2)/(nvec/sigma^2+1/lambda^2)
      pvec<-nvec*dnorm(rep(Y[i],Kval),Mvec,sqrt(Svec))
      pvec<-c(pvec,alpha*dnorm(Y[i],0,sqrt(sigma^2 + lambda^2)))/(n-1+alpha)

      pvec<-pvec/sum(pvec)
      Ksamp<-sample(c(1:(Kval+1)),size=1,prob=pvec)
      Cvec[i]<-Ksamp
      if(Ksamp == Kval+1){
        Kval<-Kval+1

```

```

        Ysum.Vec<-c(sum.Vec,Y[i])
        Nvec<-c(nvec,1)
    }else{
        Ysum.Vec<-sum.Vec
        Ysum.Vec[Ksamp]<-sum.Vec[Ksamp]+Y[i]
        Nvec<-nvec
        Nvec[Ksamp]<-nvec[Ksamp]+1
    }
}

Nsamp[irep]<-Kval

if(irep %% 500 == 0){

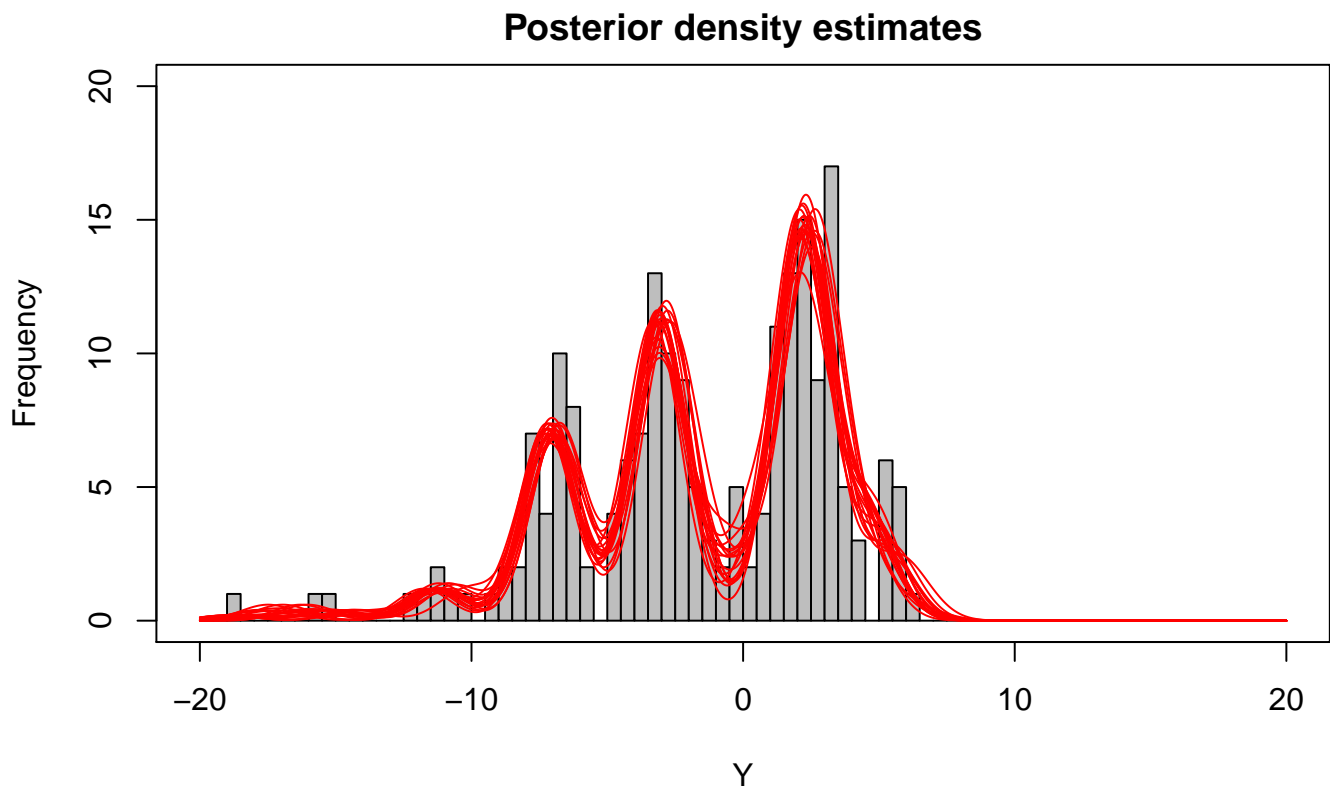
    MuVec<-(Ysum.Vec/sigma^2)/(Nvec/sigma^2+1/lambda^2)
    SigVec<-1/(Nvec/sigma^2+1/lambda^2)
    Xvec<-rnorm(Kval)*sqrt(SigVec)+MuVec
    fyv<-yv*0
    for(k in 1:Kval){
        fyv<-fyv+Nvec[k]*dnorm(yv,Xvec[k],sigma)/n
    }
    fyv.samp<-cbind(fyv.samp,fyv)
}
}

```

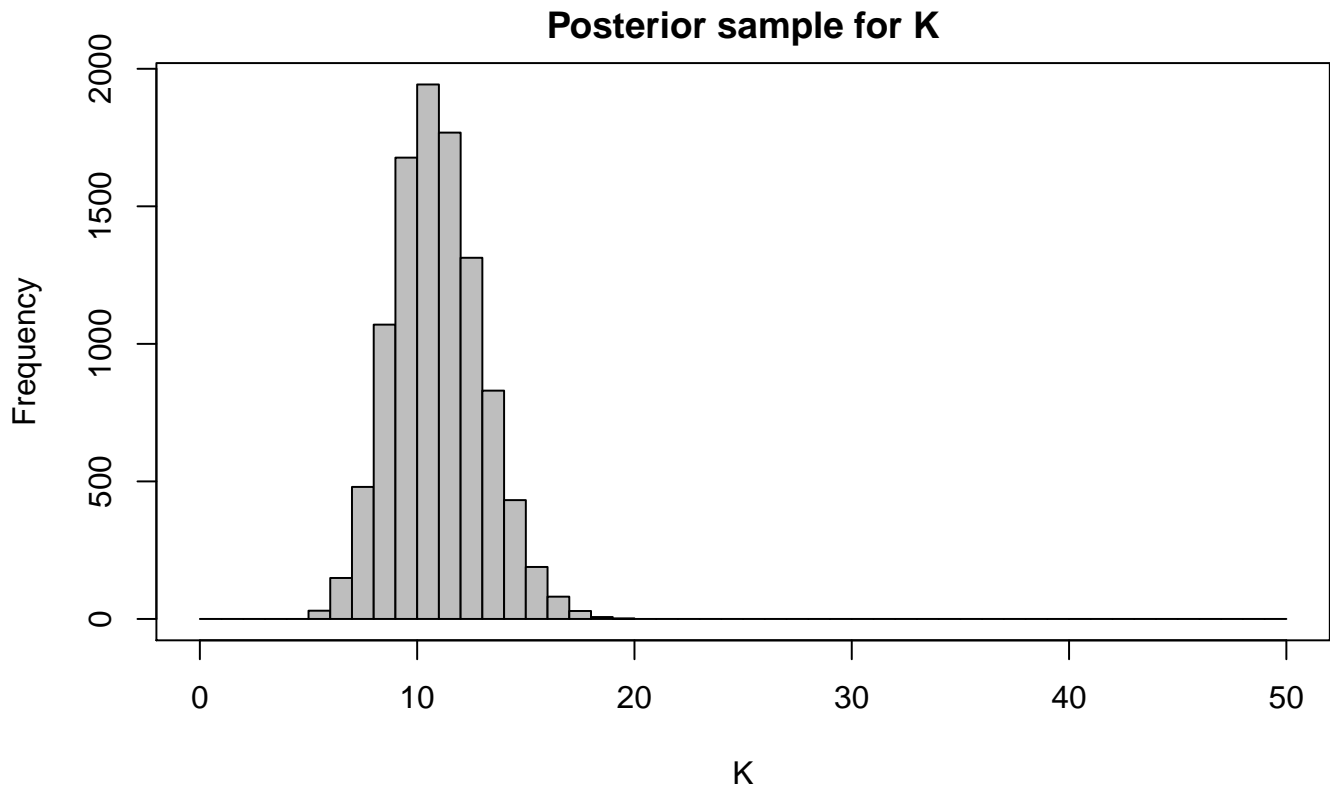
```

par(mar=c(4,4,2,0))
hist(Y,breaks=seq(-20,20,by=0.5),col='gray',ylim=range(0,20),main='Posterior density estimates');box()
for(i in 1:ncol(fyv.samp)){lines(yv,fyv.samp[,i]*n*0.5,col='red')}

```



```
hist(Nsamp,col='gray',breaks=seq(0,50,by=1),xlab='K',main='Posterior sample for K');box()
```



```
#Galaxy data using Gibbs sampler
Y.galaxy<-MASS::galaxies/10000
n.galaxy<-length(Y.galaxy)

set.seed(1010)
index<-c(1:n.galaxy)
nreps<-10000

alpha<-2.0

Pi<-Y.galaxy
Prior.K.galaxy<-numeric()

for(irep in 1:nreps){
  for(i in 1:n.galaxy){
    pvec<-rep(1/(alpha+n.galaxy-1),n.galaxy)
    pvec[i]<-alpha/(alpha+n.galaxy-1)
    ival<-sample(index,size=1,prob=pvec)
    if(ival == i){
      Pi[i]<-rnorm(1,0,lambda)
    }else{
      Pi[i]<-Pi[ival]
    }
  }

  Kval<-length(table(Pi))
  Prior.K.galaxy<-c(Prior.K.galaxy,Kval)
}

set.seed(1010)
```

```

index<-c(1:n.galaxy)
nreps<-10000
sigma<-0.1
lambda<-0.5

Pi<-Y.galaxy
Pvec<-rep(0,n.galaxy)
Posterior.K.galaxy<-numeric()
yv<-seq(0,4,by=0.01)
fyv.samp.GAL<-numeric()
for(irep in 1:nreps){

  for(i in 1:n.galaxy){
    pvec<-dnorm(Y.galaxy[i],Pi,rep(sigma,n.galaxy))
    pvec[i]<-alpha*dnorm(Y.galaxy[i],0,sqrt(sigma^2+lambda^2))
    ival<-sample(index,size=1,prob=pvec)
    if(ival == i){
      muval<-(Y.galaxy[i]/sigma^2)/(1/sigma^2+1/lambda^2)
      sval<-sqrt(1/(1/sigma^2+1/lambda^2))
      Pi[i]<-rnorm(1,muval,sval)
    }else{
      Pi[i]<-Pi[ival]
    }
  }

  Kval<-length(table(Pi))
  Posterior.K.galaxy<-c(Posterior.K.galaxy,Kval)

  if(irep %% 100 == 0){
    fyv<-yv*0
    for(k in 1:n.galaxy){
      fyv<-fyv+dnorm(yv,Pi[k],sigma)/n.galaxy
    }
    fyv.samp.GAL<-cbind(fyv.samp.GAL,fyv)
  }
}

```

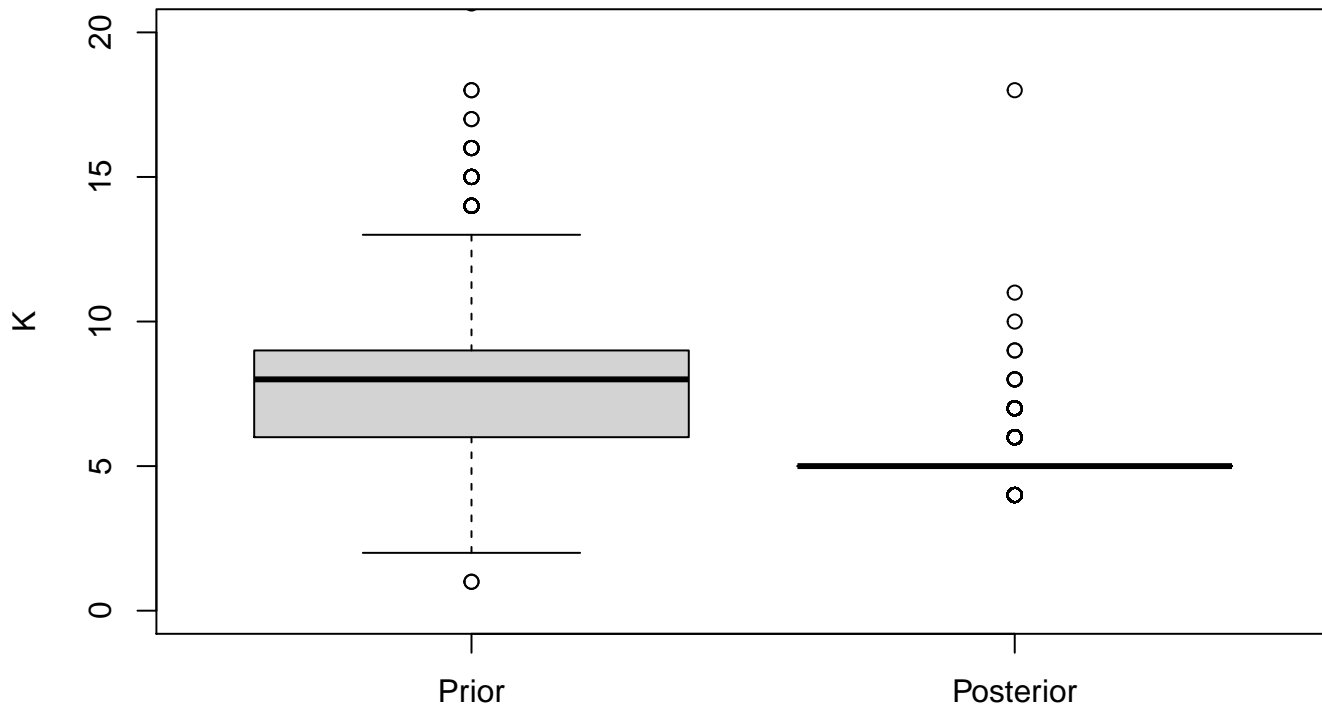
```
table(Posterior.K.galaxy)/length(Posterior.K.galaxy)
```

```
+ Posterior.K.galaxy
+   4   5   6   7   8   9   10  11  18  28
+ 0.0512 0.8120 0.1260 0.0096 0.0006 0.0002 0.0001 0.0001 0.0001 0.0001
```

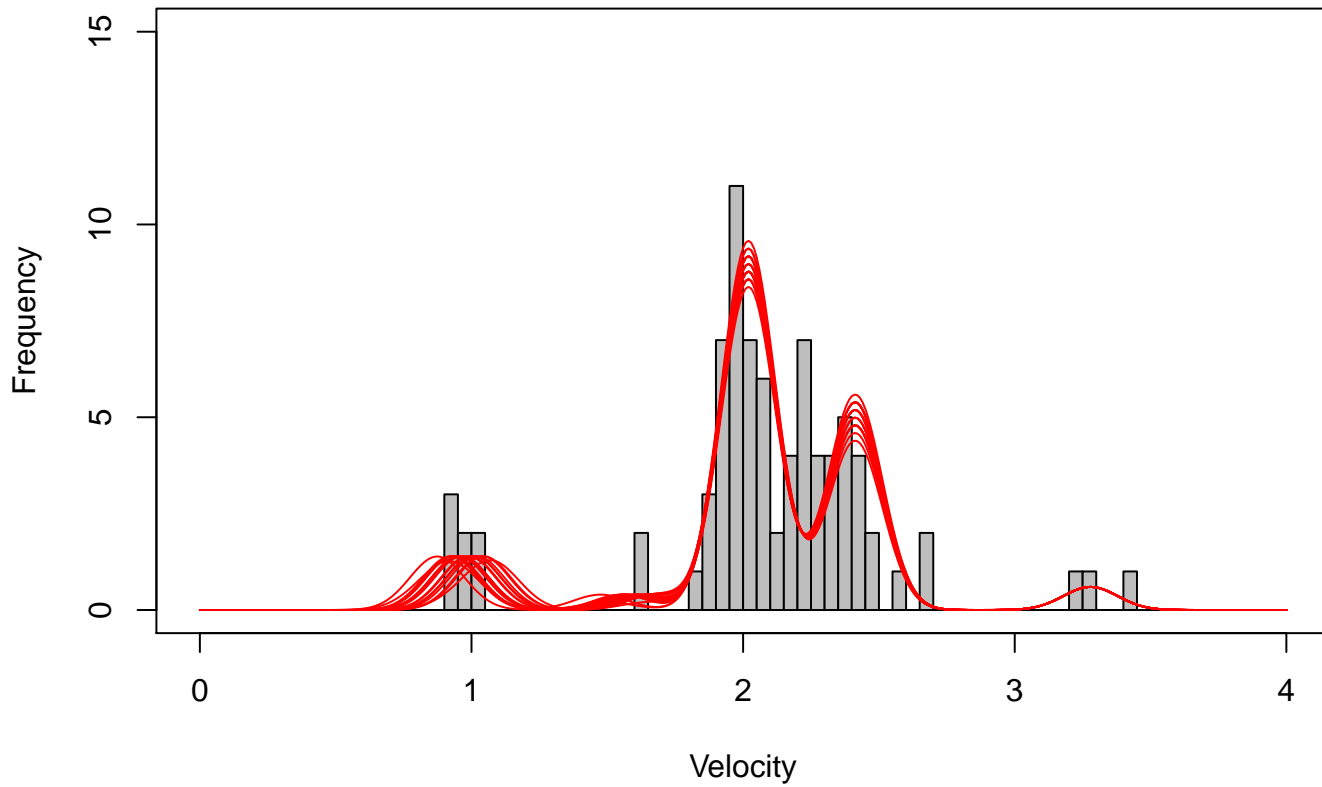
```
table(Prior.K.galaxy)/length(Prior.K.galaxy)
```

```
+ Prior.K.galaxy
+   1   2   3   4   5   6   7   8   9   10  11
+ 0.0002 0.0019 0.0135 0.0382 0.0862 0.1399 0.1638 0.1686 0.1483 0.1072 0.0651
+   12  13  14  15  16  17  18  21  28
+ 0.0383 0.0179 0.0062 0.0032 0.0009 0.0002 0.0002 0.0001 0.0001
```

### Posterior for K



### Galaxy Data



```

#Galaxy data using marginalized algorithm
Y.galaxy<-MASS::galaxies/10000
n.galaxy<-length(Y.galaxy)

set.seed(1010)
index<-c(1:n.galaxy)
nreps<-10000
alpha<-2.0
sigma<-0.1
lambda<-0.5

index<-c(1:n.galaxy)
Kval<-5
Yq<-quantile(Y.galaxy,prob=c(0:Kval)/Kval)
Cvec<-as.numeric(cut(Y.galaxy,Yq,include.lowest=T))
Ysum.Vec<-rep(0,Kval)
for(k in 1:Kval){Ysum.Vec[k]<-sum(Y.galaxy[Cvec==k])}
Nvec<-as.numeric(table(Cvec))

Posterior.K.marg<-rep(0,nreps)
yv<-seq(0,4,by=0.01)
marg.fyv.samp.GAL<-numeric()

for(irep in 1:nreps){

  for(i in 1:n.galaxy){
    sum.Vec<-Ysum.Vec
    sum.Vec[Cvec[i]]<-sum.Vec[Cvec[i]]-Y.galaxy[i]
    nvec<-Nvec
    nvec[Cvec[i]]<-nvec[Cvec[i]]-1
    if(nvec[Cvec[i]] > 0){
      mvec<-sum.Vec/nvec
      Mvec<-(sum.Vec/sigma^2)/(nvec/sigma^2+1/lambda^2)
      Svec<-sigma^2*((nvec+1)/sigma^2+1/lambda^2)/(nvec/sigma^2+1/lambda^2)
      pvec<-nvec*dnorm(rep(Y.galaxy[i],Kval),Mvec,sqrt(Svec))
      pvec<-c(pvec,alpha*dnorm(Y.galaxy[i],0,sqrt(sigma^2 + lambda^2)))
      pvec<-pvec/sum(pvec)
      Ksamp<-sample(c(1:(Kval+1)),size=1,prob=pvec)
      Cvec[i]<-Ksamp
      if(Ksamp == Kval+1){
        Kval<-Kval+1
        Ysum.Vec<-c(sum.Vec,Y.galaxy[i])
        Nvec<-c(nvec,1)
      }else{
        Ysum.Vec<-sum.Vec
        Ysum.Vec[Ksamp]<-sum.Vec[Ksamp]+Y.galaxy[i]
        Nvec<-nvec
        Nvec[Ksamp]<-nvec[Ksamp]+1
      }
    }else{
      Ktmp<-Cvec[i]
      sum.Vec<-sum.Vec[-Ktmp]
      nvec<-nvec[-Ktmp]
      Kval<-Kval-1
      Cvec[Cvec > Ktmp]<-Cvec[Cvec > Ktmp]-1

      mvec<-sum.Vec/nvec
      Mvec<-(sum.Vec/sigma^2)/(nvec/sigma^2+1/lambda^2)
      Svec<-sigma^2*((nvec+1)/sigma^2+1/lambda^2)/(nvec/sigma^2+1/lambda^2)
      pvec<-nvec*dnorm(rep(Y.galaxy[i],Kval),Mvec,sqrt(Svec))
      pvec<-c(pvec,alpha*dnorm(Y.galaxy[i],0,sqrt(sigma^2 + lambda^2)))/(n.galaxy-1+alpha

```



```

pvec<-pvec/sum(pvec)
Ksamp<-sample(c(1:(Kval+1)),size=1,prob=pvec)
Cvec[i]<-Ksamp
if(Ksamp == Kval+1){
  Kval<-Kval+1
  Ysum.Vec<-c(sum.Vec,Y.galaxy[i])
  Nvec<-c(nvec,1)
}else{
  Ysum.Vec<-sum.Vec
  Ysum.Vec[Ksamp]<-sum.Vec[Ksamp]+Y.galaxy[i]
  Nvec<-nvec
  Nvec[Ksamp]<-nvec[Ksamp]+1
}
}
}

Posterior.K.marg[irep]<-Kval

if(irep %% 100 == 0){
  MuVec<-(Ysum.Vec/sigma^2)/(Nvec/sigma^2+1/lambda^2)
  SigVec<-1/(Nvec/sigma^2+1/lambda^2)
  Xvec<-rnorm(Kval)*sqrt(SigVec)+MuVec
  fyv<-yv*0
  for(k in 1:Kval){
    fyv<-fyv+Nvec[k]*dnorm(yv,Xvec[k],sigma)/n.galaxy
  }
  marg.fyv.samp.GAL<-cbind(marg.fyv.samp.GAL,fyv)
}
}

```

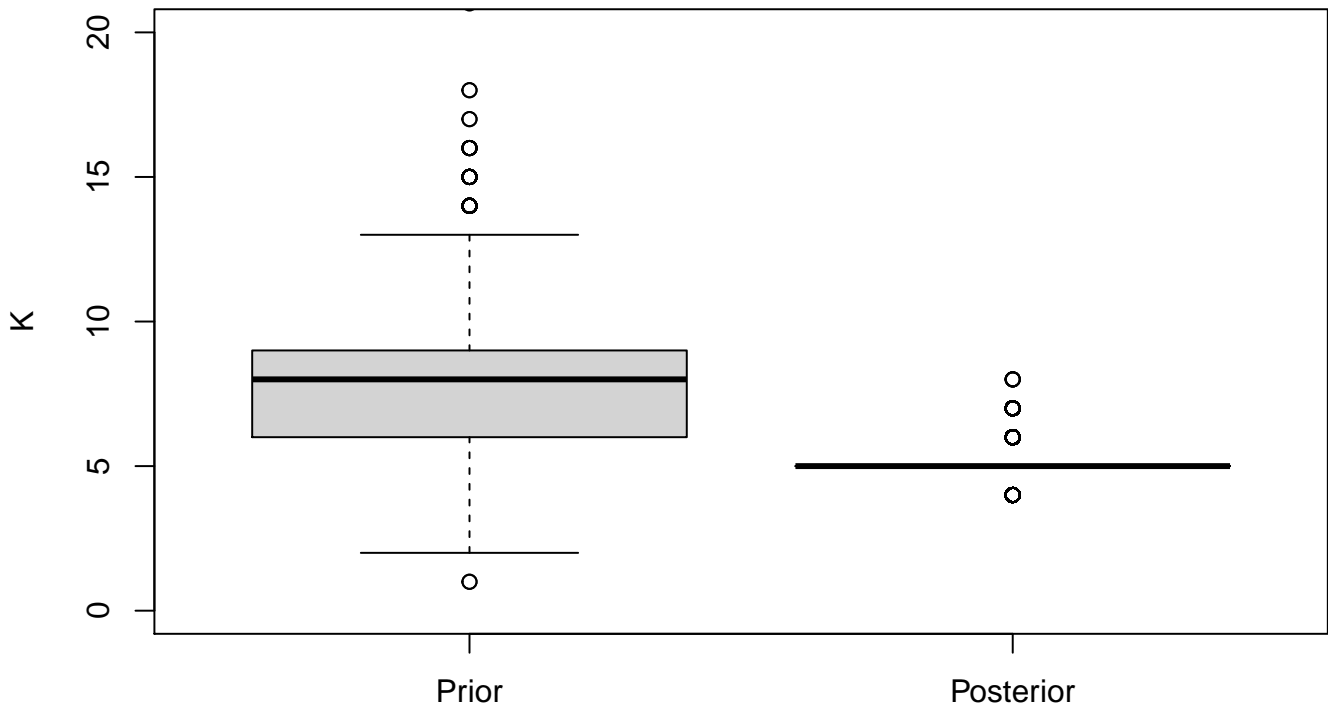
```
table(Posterior.K.marg)/nreps
```

```
+ Posterior.K.marg
+   4     5     6     7     8
+ 0.2340 0.6559 0.1021 0.0075 0.0005
```

```
table(Prior.K.galaxy)/length(Prior.K.galaxy)
```

```
+ Prior.K.galaxy
+   1     2     3     4     5     6     7     8     9    10    11
+ 0.0002 0.0019 0.0135 0.0382 0.0862 0.1399 0.1638 0.1686 0.1483 0.1072 0.0651
+   12    13    14    15    16    17    18    21    28
+ 0.0383 0.0179 0.0062 0.0032 0.0009 0.0002 0.0002 0.0001 0.0001
```

### Posterior for K



### Galaxy Data

